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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=3; day=17; hr=12; min=24; sec=12; ms=311; ]

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Application No: 10567857 Version No: 1.0

Input Set:

Output Set:

Started: 2008-03-17 10:54:35.366

Finished: 2008-03-17 10:54:36.357

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 991 ms

Total Warnings: 1

Total Errors: 0

No. of SeqIDs Defined: 4

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<141> 2008-03-17

<150> JP 2003/207397

<151> 2003-08-12

<150> JP 2004/59611

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<212> PRT

<213> Aequorea aequorea

<220>

<223> Inventor: Inouye, Satoshi

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Leu Asp Glu Met Val Tyr Lys Ala Ser Asp Ile Val Ile Asn Asn Leu  
35 40 45

Gly Ala Thr Pro Glu Gln Ala Lys Arg His Lys Asp Ala Val Glu Ala  
50 55 60

Phe Phe Gly Gly Ala Gly Met Lys Tyr Gly Val Glu Thr Asp Trp Pro  
65 70 75 80

Ala Tyr Ile Glu Gly Trp Lys Lys Leu Ala Thr Asp Glu Leu Glu Lys  
85 90 95

Tyr Ala Lys Asn Glu Pro Thr Leu Ile Arg Ile Trp Gly Asp Ala Leu  
100 105 110

Phe Asp Ile Val Asp Lys Asp Gln Asn Gly Ala Ile Thr Leu Asp Glu  
115 120 125

Trp Lys Ala Tyr Thr Lys Ala Ala Gly Ile Ile Gln Ser Ser Glu Asp  
130 135 140

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35 40 45

Asp Ile Cys Ala Lys Leu Glu Ala Thr Pro Glu Gln Thr Lys Arg His  
50 55 60

Gln Val Cys Val Glu Ala Phe Phe Arg Gly Cys Gly Met Glu Tyr Gly  
65 70 75 80

Lys Glu Ile Ala Phe Pro Gln Phe Leu Asp Gly Trp Lys Gln Leu Ala  
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Thr Ser Glu Leu Lys Lys Trp Ala Arg Asn Glu Pro Thr Leu Ile Arg  
100 105 110

Glu Trp Gly Asp Ala Val Phe Asp Ile Phe Asp Lys Asp Gly Ser Gly  
115 120 125

Thr Ile Thr Leu Asp Glu Trp Lys Ala Tyr Gly Lys Ile Ser Gly Ile  
130 135 140

Ser Pro Ser Gln Glu Asp Cys Glu Ala Thr Phe Arg His Cys Asp Leu  
145 150 155 160

Asp Asn Ser Gly Asp Leu Asp Val Asp Glu Met Thr Arg Gln His Leu  
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35 40 45

Ala Ser Asp Asp Ile Cys Ala Lys Leu Gly Ala Thr Pro Glu Gln Thr  
50 55 60

Lys Arg His Gln Asp Ala Val Glu Ala Phe Phe Lys Lys Ile Gly Met  
65 70 75 80

Asp Tyr Gly Lys Glu Val Glu Phe Pro Ala Phe Val Asp Gly Trp Lys  
85 90 95

Glu Leu Ala Asn Tyr Asp Leu Lys Leu Trp Ser Gln Asn Lys Lys Ser  
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Leu Ile Arg Asp Trp Gly Glu Ala Val Phe Asp Ile Phe Asp Lys Asp  
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130 135 140

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Cys Asp Leu Asp Asn Ser Gly Lys Leu Asp Val Asp Glu Met Thr Arg  
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Ser Asn Ile Ile Cys Lys Lys Leu Gly Ala Thr Glu Glu Gln Thr Lys  
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Leu Ala Lys Thr Glu Leu Glu Arg His Ser Lys Asn Gln Val Thr Leu  
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Ile Arg Leu Trp Gly Asp Ala Leu Phe Asp Ile Ile Asp Lys Asp Arg  
115 120 125

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Gly Ile Gln Gln Ser Arg Gly Gln Cys Glu Ala Thr Phe Ala His Cys  
145 150 155 160

Asp Leu Asp Gly Asp Gly Lys Leu Asp Val Asp Glu Met Thr Arg Gln  
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Gly Gly Ala Val Pro Tyr  
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